SEQUENCE LISTING

(1) GENERAL INFORMATION:

- (i) APPLICANT: Gunzburg, Walter H. Saller, Robert M. Salmons, Brian
- (ii) TITLE OF INVENTION:

THE USE OF THE WAP OR MMTV REGULATORY SEQUENCES FOR TARGETED EXPRESSION OF LINKED HETEROLOGOUS GENES IN HUMAN MAMMARY CELLS, INCLUDING HUMAN MAMMARY CARCINOMA CELLS

- (iii) NUMBER OF SEQUENCES: 9
 - (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C.
 - (B) STREET: Two Militia Drive
 - (C) CITY: Lexington
 - (D) STATE: Massachusetts
 - (E) COUNTRY: USA
 - (F) ZIP: 02173
 - (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk

 - (B) COMPUTER: IBM PC compatible (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
- (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER: US 09/035,596
 - (B) FILING DATE: 05-MAR-1998
 - (C) CLASSIFICATION:
- (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: PCT/EP96/03922
 (B) FILING DATE: 06-SEP-1996
- (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: DK 0976/95
 - (B) FILING DATE: 06-SEP-1995
- (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Brook, David E.

 - (B) REGISTRATION NUMBER: 22,592
 (C) REFERENCE/DOCKET NUMBER: GSF98-01A
 - (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: (781) 861-6240
 - (B) TELEFAX: (781) 861-9540
- (2) INFORMATION FOR SEQ ID NO:1:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 41 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear

| | (ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "DNA" (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1: | | | | | | | |
|------|--|----|--|--|--|--|--|--|
| ATGA | AAAGACC CCGCGGACGC GTGCGCCAGT CCTCCGATTG A | 41 | | | | | | |
| (2) | 2) INFORMATION FOR SEQ ID NO:2: | | | | | | | |
| | (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear | | | | | | | |
| | <pre>(ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "DNA"</pre> | | | | | | | |
| | (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2: | | | | | | | |
| TCAA | TCAATCACTC AGAGGAGACC | | | | | | | |
| (2) | 2) INFORMATION FOR SEQ ID NO:3: | | | | | | | |
| | (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 41 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear | | | | | | | |
| | <pre>(ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "DNA"</pre> | | | | | | | |
| | (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3: | | | | | | | |
| ACAT | CGATAG AAAAAGGGGG GAATGAAAGA CCCCGCGGAC G | 41 | | | | | | |
| (2) | INFORMATION FOR SEQ ID NO:4: | | | | | | | |
| | (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear | | | | | | | |
| | (ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "DNA" | | | | | | | |

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

ATACCGCGGA AAAAGGGGGA AATGCC

| (2) | INFORMATION FOR SEQ ID NO:5: | |
|--------------|--|----|
| | (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 33 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear | |
| | <pre>(ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "DNA"</pre> | |
| | (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5: | |
| GCGC | CGACGC GTAAGTTTAC TCAAAAAATC AGC | 33 |
| (2) | INFORMATION FOR SEQ ID NO:6: | |
| | (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 28 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear | |
| | <pre>(ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "DNA"</pre> | |
| | (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6: | |
| <u>አ</u> አሮር | GCGGCC AGGAGAAGTC ACCCTCAG | 28 |
| | INFORMATION FOR SEQ ID NO:7: | |
| (2) | (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear | |
| | <pre>(ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "DNA"</pre> | |
| | (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7: | |
| ACAC | GCGTCT GCCTCTCCCC TCAGGACACA | 30 |
| (2) | INFORMATION FOR SEQ ID NO:8: | |
| | (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear | |

| (ii) | MOLE | CULE | TYPE: | otl | ner | nuc | $:$ l ϵ | ic | aci | .d |
|------|------|------|---------|------|-----|-----|------------------|-----|-------|----|
| | (A) | DESC | CRIPTIO | : ИС | /de | esc | = | "DI | 1A '' | |
| | | | | | | | | | | |

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:

AAGCCGGATC CCTGGAGAGC ATGCAC

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- (2) INFORMATION FOR SEQ ID NO:9:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 20 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "DNA"
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:9: CGATTAGGAT CCCTGCCTCA

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